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(75) Abstract: Disclosed is a method for detecting a Chinese diabetic subject suffering from, at risk for developing, or suspected of suffering from a nephropathy. The method includes determining whether a sample from the group consisting of polymorphic sequences an I/D genotype of an ACE gene, an M235T genotype of an ACE gene, an M235T genotype of an ACR gene, an M235T genotype of an ACR gene in the promoter region, a G-308A genotype of TNP-G gene, and a complement thereof, provided that the ALR2 gene cannot be used along in which the polymorphic sequences calculated from the group consisting of polymorphic sequences an I/D genotype of an ACE gene, an M235T genotype of an ACR2 gene in the promoter region, a G-308A genotype of an ALR2 gene, and complement thereof, provided that the ALR2 gene cannot be used along in which the polymorphic sequences indicated the complement thereof, provided that the ALR2 gene cannot be used along in which the polymorphic sequences and I/D genotype of an ACR2 gene, and a complement thereof, provided that the ALR2 gene cannot be used along in which the polymorphic sequences and I/D genotype of an ACR2 gene in the promoter region, a G-308A genotype of an ACR2 gene and a complement thereof. AGT gene, a (CA) n-5' (z-2) genotype of an ALR2 gene, an C106T genotype of an ALR2 gene in the promoter region, a G-308A genotype of a TNF-α gene, and a complement thereof, provided that the ALR2 gene cannot be used alone, in which the presence of the polymorphic sequence indicates the subject suffering from, or at risk for suffering from a nephropathy. Also provided is an array for detecting a Chinese diabetic subject suffering from, or at risk for suffering from a nephropathy.



